SEQUENCE LISTING

<110> Minerva Biotechnologies Corporation

<120> Techniques and Compositions for the Diagnosis and Treatment of Cancer (MUC1)

<130> M1015.70089W000

<140> not yet assigned

<141> 2004-08-26

<150> US 60/498,260

<151> 2003-08-26

<160> 66

<170> PatentIn version 3.3

<210> 1

<211> 39

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Peptide

<400> 1

Gly Thr Ile Asn Val His Asp Val Glu Thr Gln Phe Asn Gln Tyr Lys 1 5 10 15

Thr Glu Ala Ala Ser Pro Tyr Asn Leu Thr Ile Ser Asp Val Ser Val 20 25 30

Ser His His His His His 35

<210> 2

<211> 51

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Peptide

<400> 2

Gly Thr Ile Asn Val His Asp Val Glu Thr Gln Phe Asn Gln Tyr Lys
1 5 10 15

Thr Glu Ala Ala Ser Pro Tyr Asn Leu Thr Ile Ser Asp Val Ser Val 20 25 30

Ser Asp Val Pro Phe Pro Phe Ser Ala Gln Ser Gly Ala His His His 35 40 45

PCT/US2004/027954 WO 2005/019269

His His His 50

<210> 3

<211> 54 <212> PRT <213> Artificial Sequence

<220>

<223> Synthetic Peptide

<400> 3

Val Gln Leu Thr Leu Ala Phe Arg Glu Gly Thr Ile Asn Val His Asp

Val Glu Thr Gln Phe Asn Gln Tyr Lys Thr Glu Ala Ala Ser Pro Tyr

Asn Leu Thr Ile Ser Asp Val Ser Val Ser Asp Val Pro Phe Pro Phe

His His His His His 50

<210> 4 <211> 31 <212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Peptide

<400> 4

His His His His His Gly Phe Leu Gly Leu Ser Asn Ile Lys Phe 10

Arg Pro Gly Ser Val Val Gln Leu Thr Leu Ala Phe Arg Glu

<210> 5 <211> 46 <212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Peptide

<400> 5

Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His Gly

Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala Pro

Pro Ala His Gly Val Thr Ser Ala His His His His His

<210> 6

<211> 33 <212> PRT <213> Artificial Sequence

<220>

<223> Synthetic Peptide

<400> 6

Gly Thr Ile Asn Val His Asp Val Glu Thr Gln Phe Asn Gln Tyr Lys

Thr Glu Ala Ala Ser Pro Tyr Asn Leu Thr Ile Ser Asp Val Ser Val

<210> 7 <211> 45 <212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Peptide

<400> 7

Gly Thr Ile Asn Val His Asp Val Glu Thr Gln Phe Asn Gln Tyr Lys

Thr Glu Ala Ala Ser Pro Tyr Asn Leu Thr Ile Ser Asp Val Ser Val

Ser Asp Val Pro Phe Pro Phe Ser Ala Gln Ser Gly Ala

<210> 8

<211> 25

<212> PRT

<213> Homo sapiens

<400> 8

Gly Phe Leu Gly Leu Ser Asn Ile Lys Phe Arg Pro Gly Ser Val Val

Val Gln Leu Thr Leu Ala Phe Arg Glu

WO 2005/019269 PCT/US2004/027954

<210> 9

<211> 40

<212> PRT

<213> Homo sapiens

<400> 9

Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His Gly
1 5 10 15

Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala Pro
20 25 30

Pro Ala His Gly Val Thr Ser Ala 35 40

<210> 10

<211> 1255

<212> PRT

<213> Homo sapiens

<400> 10

Met Thr Pro Gly Thr Gln Ser Pro Phe Phe Leu Leu Leu Leu Thr 1 5 10 15

Val Leu Thr Val Val Thr Gly Ser Gly His Ala Ser Ser Thr Pro Gly 20 25 30

Gly Glu Lys Glu Thr Ser Ala Thr Gln Arg Ser Ser Val Pro Ser Ser 35 40 45

Thr Glu Lys Asn Ala Val Ser Met Thr Ser Ser Val Leu Ser Ser His 50 55 60

Ser Pro Gly Ser Gly Ser Ser Thr Thr Gln Gly Gln Asp Val Thr Leu 65 70 75 80

Ala Pro Ala Thr Glu Pro Ala Ser Gly Ser Ala Ala Thr Trp Gly Gln 85 90 95

Asp Val Thr Ser Val Pro Val Thr Arg Pro Ala Leu Gly Ser Thr Thr 100 105 110

Pro Pro Ala His Asp Val Thr Ser Ala Pro Asp Asn Lys Pro Ala Pro 115 120 125

Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr 130 135 140

Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser 145 150 155 160

Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His 165 170 175

Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala 180 185 190

								-								
Pro	Pro	Ala 195	His	Gly	Val	Thr	Ser 200	Ala	Pro	Asp	Thr	Arg 205	Pro	Ala	Pro	
Gly	Ser 210	Thr	Ala	Pro	Pro	Ala 215	His	Gly	Val	Thr	Ser 220	Ala	Pro	Asp	Thr	
Arg 225	Pro	Ala	Pro	Gly	Ser 230	Thr	Ala	Pro	Pro	Ala 235	His	Gly	Val	Thr	Ser 240	
Ala	Pro	Asp	Thr	Arg 245	Pro	Ala	Pro	Gly	Ser 250	Thr	Ala	Pro	Pro	Ala 255	His	
Gly	Val	Thr	Ser 260	Ala	Pro	Asp	Thr	Arg 265	Pro	Ala	Pro	Gly	Ser 270	Thr	Ala	
Pro	Pro	Ala 275	His	Gly	Val	Thr	Ser 280	Ala	Pro	Asp	Thr	Arg 285	Pro	Ala	Pro	
Gly	Ser 290	Thr	Ala	Pro	Pro	Ala 295	His	Gly	Val	Thr	Ser 300	Ala	Pro	Asp	Thr	
Arg 305	Pro	Ala	Pro	Gly	Ser 310	Thr	Ala	Pro	Pro	Ala 315	His	Gly	Val	Thr	Ser 320	
Ala	Pro	Asp	Thr	Arg 325	Pro	Ala	Pro	Gly	Ser 330	Thr	Ala	Pro	Pro	Ala 335	His	
Gly	Val	Thr	Ser 340	Ala	Pro	Asp	Thr	Arg 345	Pro	Ala	Pro	Gly	Ser 350	Thr	Ala	
Pro	Pro	Ala 355	His	Gly	Val	Thr	Ser 360	Ala	Pro	Asp	Thr	Arg 365	Pro	Ala	Pro	
Gly	Ser 370	Thr	Ala	Pro	Pro	Ala 375	His	Gly	Val	Thr	Ser 380	Ala	Pro	Asp	Thr	
Arg 385	Pro	Ala	Pro	Gly	Ser 390	Thr	Ala	Pro	Pro	Ala 395	His	Gly	Val	Thr	Ser 400	
Ala	Pro	Asp	Thr	Arg 405	Pro	Ala	Pro	Glу	Ser 410	Thr	Ala	Pro	Pro	Ala 415	His	
Gly	Val	Thr	Ser 420	Ala	Pro	Asp	Thr	Arg 425	Pro	Ala	Pro	Gly	Ser 430	Thr	Ala	
Pro	Pro	Ala 435	His	Gly	Val	Thr	Ser 440	Ala	Pro	Asp	Thr	Arg 445	Pro	Ala	Pro	
Gly	Ser 450	Thr	Ala	Pro	Pro	Ala 455	His	Gly	Val	Thr	Ser 460	Ala	Pro	Asp	Thr	
Arg 465	Pro	Ala	Pro	Gly	Ser 470	Thr	Ala	Pro	Pro	Ala 475	His	Gly	Val	Thr	Ser 480	
Ala	Pro	Asp	Thr	Arg 485	Pro	Ala	Pro	Gly	Ser 490	Thr	Ala	Pro	Pro	Ala 495	His	
Gly	Val	Thr	Ser 500	Ala	Pro	Asp	Thr	Arg 505	Pro	Ala	Pro	Gly	Ser 510	Thr	Ala	

WO 2005/019269 PCT/US2004/027954 6/32

Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr 535 Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala 585 Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr 615 Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro 680 Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His 730 Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His 810 Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala 820 825

Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro 835 840 845

- Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr 850 855 860
- Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser 865 870 875 880
- Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His 885 890 895
- Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala 900 905 910
- Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro 915 920 925
- Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Asn 930 940
- Arg Pro Ala Leu Gly Ser Thr Ala Pro Pro Val His Asn Val Thr Ser 945 950 955 960
- Ala Ser Gly Ser Ala Ser Gly Ser Ala Ser Thr Leu Val His Asn Gly 965 970 975
- Thr Ser Ala Arg Ala Thr Thr Thr Pro Ala Ser Lys Ser Thr Pro Phe 980 985 990
- Ser Ile Pro Ser His His Ser Asp Thr Pro Thr Thr Leu Ala Ser His 995 1000 1005
- Ser Thr Lys Thr Asp Ala Ser Ser Thr His His Ser Ser Val Pro 1010 1015 1020
- Pro Leu Thr Ser Ser Asn His Ser Thr Ser Pro Gln Leu Ser Thr 1025
- Gly Val Ser Phe Phe Phe Leu Ser Phe His Ile Ser Asn Leu Gln 1040 1045 1050
- Phe Asn Ser Ser Leu Glu Asp Pro Ser Thr Asp Tyr Tyr Gln Glu 1055 1060 1065
- Leu Gln Arg Asp Ile Ser Glu Met Phe Leu Gln Ile Tyr Lys Gln 1070 1075 1080
- Gly Gly Phe Leu Gly Leu Ser Asn Ile Lys Phe Arg Pro Gly Ser 1085 1090 1095
- Val Val Gln Leu Thr Leu Ala Phe Arg Glu Gly Thr Ile Asn 1100 1105 1110
- Val His Asp Val Glu Thr Gln Phe Asn Gln Tyr Lys Thr Glu Ala 1115 1120 1125
- Ala Ser Arg Tyr Asn Leu Thr Ile Ser Asp Val Ser Val Ser Asp 1130 1135 1140

Val Pro Phe Pro Phe Ser Ala Gln Ser Gly Ala Gly Val Pro Gly 1145 1150

Trp Gly Ile Ala Leu Leu Val Leu Val Cys Val Leu Val Ala Leu 1165

Ala Ile Val Tyr Leu Ile Ala Leu Ala Val Cys Gln Cys Arg Arg 1180

Lys Asn Tyr Gly Gln Leu Asp Ile Phe Pro Ala Arg Asp Thr Tyr 1190 1195

His Pro Met Ser Glu Tyr Pro Thr Tyr His Thr His Gly Arg Tyr 1210

Val Pro Pro Ser Ser Thr Asp Arg Ser Pro Tyr Glu Lys Val Ser 1225

Ala Gly Asn Gly Gly Ser Ser Leu Ser Tyr Thr Asn Pro Ala Val 1240

Ala Ala Ser Ala Asn Leu 1250 1255

<210> 11

<211> 302 <212> PRT <213> Homo sapiens

<400> 11

Ala Ala Ala Lys Glu Gly Lys Lys Ser Arg Asp Arg Glu Arg Pro Pro

Ser Val Pro Ala Leu Arg Glu Gln Pro Pro Glu Thr Glu Pro Gln Pro

Ala Trp Lys Met Pro Arg Ser Cys Cys Ser Arg Ser Gly Ala Leu Leu 40

Leu Ala Leu Leu Gln Ala Ser Met Glu Val Arg Gly Trp Cys Leu

Glu Ser Ser Gln Cys Gln Asp Leu Thr Thr Glu Ser Asn Leu Leu Glu

Cys Ile Arg Ala Cys Lys Pro Asp Leu Ser Ala Glu Thr Pro Met Phe

Pro Gly Asn Gly Asp Glu Gln Pro Leu Thr Glu Asn Pro Arg Lys Tyr

Val Met Gly His Phe Arg Trp Asp Arg Phe Gly Arg Arg Asn Ser Ser

Ser Ser Gly Ser Ser Gly Ala Gly Gln Lys Arg Glu Asp Val Ser Ala 135

Gly Glu Asp Cys Gly Pro Leu Pro Glu Gly Gly Pro Glu Pro Arg Ser

Asp Gly Ala Lys Pro Gly Pro Arg Glu Gly Lys Arg Ser Tyr Ser Met 165 170 175

Glu His Phe Arg Trp Gly Lys Pro Val Gly Lys Lys Arg Arg Pro Val 180 185 190

Lys Val Tyr Pro Asn Gly Ala Glu Asp Glu Ser Ala Glu Ala Phe Pro 195 200 205

Leu Glu Phe Lys Arg Glu Leu Thr Gly Gln Arg Leu Arg Glu Gly Asp 210 215 220

Gly Pro Asp Gly Pro Ala Asp Asp Gly Ala Gly Ala Gln Ala Asp Leu 225 230 235 240

Glu His Ser Leu Leu Val Ala Ala Glu Lys Lys Asp Glu Gly Pro Tyr 245 250 255

Arg Met Glu His Phe Arg Trp Gly Ser Pro Pro Lys Asp Lys Arg Tyr 260 265 270

Gly Gly Phe Met Thr Ser Glu Lys Ser Gln Thr Pro Leu Val Thr Leu 275 280 285

Phe Lys Asn Ala Ile Ile Lys Asn Ala Tyr Lys Lys Gly Glu 290 295 300

<210> 12

<211> 31

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Peptide

<400> 12

His His His His His Ser Ser Ser Gly Ser Ser Ser Gly 1 5 10 15

Ser Ser Ser Gly Gly Arg Gly Asp Ser Gly Arg Gly Asp Ser 20 25 30

<210> 13

<211> 19

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Peptide

<400> 13

His His His His His Arg Gly Glu Phe Thr Gly Thr Tyr Ile Thr 1 5 10 15

10/32

Ala Val Thr

```
<210> 14
<211> 12
<212> PRT
<213> Homo sapiens
<400> 14
Thr Phe Ile Ala Ile Lys Pro Asp Gly Val Gln Arg
1 5
<210> 15
<211> 18
<212> PRT
<213> Homo sapiens
<220>
<221> misc_feature <222> (3)..(3)
<223> Xaa can be any naturally occurring amino acid
<400> 15
Val Met Xaa Leu Gly Glu Thr Asn Pro Ala Asp Ser Lys Pro Gly Thr
        5
                                      10 , 15
Ile Arg
<210> 16
<211> 17
<212> PRT
<213> Homo sapiens
<400> 16
Val Met Leu Gly Glu Thr Asn Pro Ala Asp Ser Lys Pro Gly Thr Ile
Arg
<210> 17
<211> 10
<212> PRT
<213> Homo sapiens
<400> 17
Asn Ile Ile His Gly Ser Asp Ser Val Lys
```

```
<210> 18
<211> 9
<212> PRT
<213> Homo sapiens
<400> 18
Gly Leu Val Gly Glu Ile Ile Lys Arg
1 5
<210> 19
<211> 8
<212> PRT
<213> Homo sapiens
<400> 19
Gly Leu Val Gly Glu Ile Ile Lys
<210> 20
<211> 21
<212> PRT
<213> Homo sapiens
<220>
<221> misc_feature
<222> (3)..(3) <223> Xaa can be any naturally occurring amino acid
<220>
<221> misc_feature
<222> (12)..(12)
<223> Xaa can be any naturally occurring amino acid
<400> 20
Tyr Met Xaa His Ser Gly Pro Val Val Ala Met Xaa Val Trp Glu Gly
Leu Asn Val Val Lys
              20
<210> 21
<211> 19
<212> PRT
<213> Homo sapiens
<400> 21
Ala Ala Phe Asp Asp Ala Ile Ala Glu Leu Asp Thr Leu Ser Glu Glu
                                         10
Ser Tyr Lys
```

WO 2005/019269 PCT/US2004/027954

```
<210> 22
<211> 18
<212> PRT
<213> Homo sapiens
<220>
<221> misc_feature
<222> (8)..(8) <223> Xaa can be any naturally occurring amino acid
<400> 22
Ala Ala Ser Asp Ile Ala Met Xaa Thr Glu Leu Pro Pro Thr His Pro
Ile Arg
<210> 23
<211> 11
<212> PRT
<213> Homo sapiens
<400> 23
Tyr Leu Ala Glu Phe Ala Thr Gly Asn Asp Arg
<210> 24
<211> 10
<211> 10
<212> PRT
<213> Homo sapiens
 <400> 24
Asp Ser Thr Leu Ile Met Gln Leu Leu Arg
 <210> 25
 <211> 9
 <212> PRT
<213> Homo sapiens
 <400> 25
 Tyr Asp Glu Met Val Glu Ser Met Lys
 1 5
 <210> 26
 <211> 14
<212> PRT
<213> Homo sapiens
 <220>
 <221> misc_feature
```

13/32

```
<222> (5)..(5) 
<223> Xaa can be any naturally occurring amino acid
<400> 26
Val Ala Gly Met Xaa Asp Val Glu Leu Thr Val Glu Glu Arg
<210> 27
<211> 12
<212> PRT
<213> Homo sapiens
<400> 27
His Leu Ile Pro Ala Ala Asn Thr Gly Glu Ser Lys
<210> 28
<211> 19
<212> PRT
<213> Homo sapiens
<220>
<221> misc_feature
<222> (12)..(12)
<223> Xaa can be any naturally occurring amino acid
<400> 28
Asp Pro Asp Ala Gln Pro Gly Gly Glu Leu Met Xaa Leu Gly Gly Thr
                                                                15
Asp Ser Lys
<210> 29
<211> 18
<212> PRT
<213> Homo sapiens
<400> 29
Asp Pro Asp Ala Gln Pro Gly Gly Glu Leu Met Leu Gly Gly Thr Asp
Ser Lys
<210> 30
<211> 18
<212> PRT
<213> Homo sapiens
<220>
<221> misc_feature
<222> (15)..(15)
```

14/32

```
<223> Xaa can be any naturally occurring amino acid
<400> 30
Ile Ser Val Asn Asn Val Leu Pro Val Phe Asp Asn Leu Met Xaa Gln
Gln Lys
<210> 31
<211> 17
<212> PRT
<213> Homo sapiens
<400> 31
Ile Ser Val Asn Asn Val Leu Pro Val Phe Asp Asn Leu Met Gln Gln
                                       10
Lys
<210> 32
<211> 10
<212> PRT
<213> Homo sapiens
<400> 32
Gln Pro Gly Ile Thr Phe Ile Ala Ala Lys
<210> 33
<211> 16
<212> PRT
<213> Homo sapiens
<400> 33
Gly Leu Gly Thr Asp Glu Glu Ser Ile Leu Thr Leu Leu Thr Ser Arg
                 5
<210> 34
<211> 13
<212> PRT
<213> Homo sapiens
<400> 34
Asp Leu Leu Asp Asp Leu Lys Ser Glu Leu Thr Gly Lys
                5
<210> 35
<211> 9
<212> PRT
```

<213> Homo sapiens

<400> 35

Ser Glu Ile Asp Leu Phe Asn Ile Arg

<210> 36 <211> 45 <212> PRT

<213> Homo sapiens

<400> 36

Gly Thr Ile Asn Val His Asp Val Glu Thr Gln Phe Asn Gln Tyr Lys

Thr Glu Ala Ala Ser Arg Tyr Asn Leu Thr Ile Ser Asp Val Ser Val

Ser Asp Val Pro Phe Pro Phe Ser Ala Gln Ser Gly Ala

<210> 37

<211> 146 <212> PRT <213> Homo sapiens

<400> 37

Gly Thr Ile Asn Val His Asp Val Glu Thr Gln Phe Asn Gln Tyr Lys

Thr Glu Ala Ala Ser Arg Tyr Asn Leu Thr Ile Ser Asp Val Ser Val

Ser Asp Val Pro Phe Pro Phe Ser Ala Gln Ser Gly Ala Gly Val Pro 40

Gly Trp Gly Ile Ala Leu Leu Val Leu Val Cys Val Leu Val Ala Leu

Ala Ile Val Tyr Leu Ile Ala Leu Ala Val Cys Gln Cys Arg Arg Lys

Asn Tyr Gly Gln Leu Asp Ile Phe Pro Ala Arg Asp Thr Tyr His Pro

Met Ser Glu Tyr Pro Thr Tyr His Thr His Gly Arg Tyr Val Pro Pro

Ser Ser Thr Asp Arg Ser Pro Tyr Glu Lys Val Ser Ala Gly Asn Gly

Gly Ser Ser Leu Ser Tyr Thr Asn Pro Ala Val Ala Ala Ala Ser Ala 135 140

Asn Leu

145

<210> 38 <211> 171 <212> PRT

<213> Homo sapiens

<400> 38

Gly Phe Leu Gly Leu Ser Asn Ile Lys Phe Arg Pro Gly Ser Val Val 1 5 10 15

Val Gln Leu Thr Leu Ala Phe Arg Glu Gly Thr Ile Asn Val His Asp 20 25 30

Val Glu Thr Gln Phe Asn Gln Tyr Lys Thr Glu Ala Ala Ser Arg Tyr 35 40 45

Asn Leu Thr Ile Ser Asp Val Ser Val Ser Asp Val Pro Phe Fro Phe 50 55 60

Ser Ala Gln Ser Gly Ala Gly Val Pro Gly Trp Gly Ile Ala Leu Leu 65 70 75 80

Val Leu Val Cys Val Leu Val Ala Leu Ala Ile Val Tyr Leu Ile Ala 85 90 95

Leu Ala Val Cys Gln Cys Arg Arg Lys Asn Tyr Gly Gln Leu Asp Ile 100 105 110

Phe Pro Ala Arg Asp Thr Tyr His Pro Met Ser Glu Tyr Pro Thr Tyr 115 120 125

His Thr His Gly Arg Tyr Val Pro Pro Ser Ser Thr Asp Arg Ser Pro 130 135 140

Tyr Glu Lys Val Ser Ala Gly Asn Gly Gly Ser Ser Leu Ser Tyr Thr 145 150 155 160

Asn Pro Ala Val Ala Ala Ala Ser Ala Asn Leu 165 170

<210> 39

<211> 275

<212> PRT

<213> Homo sapiens

<400> 39

Ala Thr Thr Thr Pro Ala Ser Lys Ser Thr Pro Phe Ser Ile Pro Ser 1 5 10 15

His His Ser Asp Thr Pro Thr Thr Leu Ala Ser His Ser Thr Lys Thr 20 25 30

Asp Ala Ser Ser Thr His His Ser Thr Val Pro Pro Leu Thr Ser Ser 35 40 45

Asn His Ser Thr Ser Pro Gln Leu Ser Thr Gly Val Ser Phe Phe 50 60

Leu Ser Phe His Ile Ser Asn Leu Gln Phe Asn Ser Ser Leu Glu Asp 65 70 75 80

Pro Ser Thr Asp Tyr Tyr Gln Glu Leu Gln Arg Asp Ile Ser Glu Met 85 90 95

Phe Leu Gln Ile Tyr Lys Gln Gly Gly Phe Leu Gly Leu Ser Asn Ile 100 105 110

Lys Phe Arg Pro Gly Ser Val Val Val Gln Leu Thr Leu Ala Phe Arg 115 120 125

Glu Gly Thr Ile Asn Val His Asp Val Glu Thr Gln Phe Asn Gln Tyr 130 135 140

Lys Thr Glu Ala Ala Ser Arg Tyr Asn Leu Thr Ile Ser Asp Val Ser 145 150 155 160

Val Ser Asp Val Pro Phe Pro Phe Ser Ala Gln Ser Gly Ala Gly Val 165 170 175

Pro Gly Trp Gly Ile Ala Leu Leu Val Leu Val Cys Val Leu Val Ala 180 185 190

Leu Ala Ile Val Tyr Leu Ile Ala Leu Ala Val Cys Gln Cys Arg Arg 195 200 205

Lys Asn Tyr Gly Gln Leu Asp Ile Phe Pro Ala Arg Asp Thr Tyr His 210 215 220

Pro Met Ser Glu Tyr Pro Thr Tyr His Thr His Gly Arg Tyr Val Pro 225 230 235 240

Pro Ser Ser Thr Asp Arg Ser Pro Tyr Glu Lys Val Ser Ala Gly Asn 245 250 255

Gly Gly Ser Ser Leu Ser Tyr Thr Asn Pro Ala Val Ala Ala Ala Ser 260 265 270

Ala Asn Leu 275

<210> 40

<211> 233

<212> PRT

<213> Homo sapiens

<400> 40

Gly Ser Gly His Ala Ser Ser Thr Pro Gly Gly Glu Lys Glu Thr Ser

10 15

Ala Thr Gln Arg Ser Ser Val Pro Ser Ser Thr Glu Lys Asn Ala Phe 20 25 30

Asn Ser Ser Leu Glu Asp Pro Ser Thr Asp Tyr Tyr Gln Glu Leu Gln 35

WO 2005/019269 PCT/US2004/027954 18/32

Arg Asp Ile Ser Glu Met Phe Leu Gln Ile Tyr Lys Gln Gly Gly Phe 50 55 60

Leu Gly Leu Ser Asn Ile Lys Phe Arg Pro Gly Ser Val Val Gln 65 70 75 80

Leu Thr Leu Ala Phe Arg Glu Gly Thr Ile Asn Val His Asp Met Glu 85 90 95

Thr Gln Phe Asn Gln Tyr Lys Thr Glu Ala Ala Ser Arg Tyr Asn Leu 100 105 110

Thr Ile Ser Asp Val Ser Val Ser Asp Val Pro Phe Pro Phe Ser Ala 115 120 125

Gln Ser Gly Ala Gly Val Pro Gly Trp Gly Ile Ala Leu Leu Val Leu 130 135 140

Val Cys Val Leu Val Ala Leu Ala Ile Val Tyr Leu Ile Ala Leu Ala 145 150 155 160

Val Cys Gln Cys Arg Arg Lys Asn Tyr Gly Gln Leu Asp Ile Phe Pro 165 170 175

Ala Arg Asp Thr Tyr His Pro Met Ser Glu Tyr Pro Thr Tyr His Thr 180 185 190

His Gly Arg Tyr Val Pro Pro Ser Ser Thr Asp Arg Ser Pro Tyr Glu 195 200 205

Lys Val Ser Ala Gly Asn Gly Gly Ser Ser Leu Ser Tyr Thr Asn Pro 210 215 220

Ala Val Ala Ala Thr Ser Ala Asn Leu 225 230

<210> 41

<211> 863

<212> PRT

<213> Homo sapiens

<400> 41

Leu Asp Pro Arg Val Arg Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro 1 5 10 15

Gly Ser Thr Ala Pro Gln Ala His Gly Val Thr Ser Ala Pro Asp Thr 20 25 30

Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His Gly Val Thr Ser 35 40 45

Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala Pro Pro Ala His 50 55 60

Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro Gly Ser Thr Ala 65 70 75 80

Pro Pro Ala His Gly Val Thr Ser Ala Pro Asp Thr Arg Pro Ala Pro 85 90 95

Gly	Ser	Thr	Ala 100	Pro	Pro	Ala	His	Gly 105	Val	Thr	Ser	Ala	Pro 110	Asp	Thr
Arg	Pro	Ala 115	Pro	Gly	Ser	Thr	Ala 120	Pro	Pro	Ala	His	Gly 125	Val	Thr	Ser
Ala	Pro 130	Asp	Thr	Arg	Pro	Ala 135	Pro	Gly	Ser	Thr	Ala 140	Pro	Pro	Ala	His
Gly 145	Val	Thr	Ser	Ala	Pro 150	Asp	Thr	Arg	Pro	Ala 155	Pro	Gly	Ser	Thr	Ala 160
Pro	Pro	Ala	His	Gly 165	Val	Thr	Ser	Ala	Pro 170	Asp	Thr	Arg	Pro	Ala 175	Pro
Gly	Ser	Thr	Ala 180	Pro	Pro	Ala	His	Gly 185	Val	Thr	Ser	Ala	Pro 190	Asp	Thr
Arg	Pro	Ala 195	Pro	Gly	Ser	Thr	Ala 200	Pro	Pro	Ala	His	Gly 205	Val	Thr	Ser
Ala	Pro 210	Asp	Thr	Arg	Pro	Ala 215	Pro	Gly	Ser	Thr	Ala 220	Pro	Pro	Ala	His
Gly 225	Val	Thr	Ser	Ala	Pro 230	Asp	Thr	Arg	Pro	Ala 235	Pro	Gly	Ser	Thr	Ala 240
Pro	Pro	Ala	His	Gly 245	Val	Thr	Ser	Ala	Pro 250	Asp	Thr	Arg	Pro	Ala 255	Pro
Gly	Ser	Thr	Ala 260	Pro	Pro	Ala	His	Gly 265	Val	Thr	Ser	Ala	Pro 270	Asp	Thr
Arg	Pro	Ala 275	Pro	Gly	Ser	Thr	Ala 280	Pro	Pro	Ala	His	Gly 285	Val	Thr	Ser
	290					295					300				His
Gly 305	Val	Thr	Ser	Ala	Pro 310	Asp	Thr	Arg	Pro	Ala 315	Pro	Gly	Ser	Thr	Ala 320
Pro	Pro	Ala	His	Gly 325	Val	Thr	Ser	Ala	Pro 330	Asp	Thr	Arg	Pro	Ala 335	Pro
Gly	Ser	Thr	Ala 340	Pro	Pro	Ala	Hìs	Gly 345		Thr	Ser	Ala	Pro 350		Thr
Arg	Pro	Ala 355		Gly	Ser	Thr	Ala 360	Pro	Pro	Ala	His	Gly 365	Val	Thr	Ser
Ala	Pro 370	Asp	Thr	Arg	Pro	Ala 375	Pro	Gly	Ser	Thr	Ala 380		Pro	Ala	His
Gly 385		Thr	Ser	Ala	Pro 390		Thr	Arg	Pro	Ala 395		Gly	Ser	Thr	Ala 400
Pro	Pro	Ala	His	Gly 405		Thr	Ser	Ala	Pro 410		Thr	Arg	Pro	Ala 415	Pro

	Gly	Ser	Thr	Ala 420	Pro	Pro	Ala	His	Gly 425	Val	Thr	Ser	Ala	Pro 430	Asp	Thr
	Arg	Pro	Ala 435	Pro	Gly	Ser	Thr	Ala 440	Pro	Pro	Ala	His	Gly 445	Val	Thr	Ser
:	Ala	Pro 450	Asp	Thr	Arg	Pro	Ala 455	Pro	Gly	Ser	Thr	Ala 460	Pro	Pro	Ala	His
,	Gly 465	Val	Thr	Ser	Ala	Pro 470	Asp	Thr	Arg	Pro	Ala 475	Pro	Gly	Ser	Thr	Ala 480
	Pro	Pro	Ala	His	Gly 485	Val	Thr	Ser	Ala	Pro 490	Asp	Thr	Arg	Pro	Ala 495	Pro
	Glу	Ser	Thr	Ala 500	Pro	Pro	Ala	His	Gly 505	Val	Thr	Ser	Ala	Pro 510	Asp	Thr
	Arg	Pro	Ala 515	Pro	Gly	Ser	Thr	Ala 520	Pro	Pro	Ala	His	Gly 525	Val	Thr	Ser
	Ala	Pro 530	Asp	Thr	Arg	Pro	Ala 535	Pro	Gly	Ser	Thr	Ala 540	Pro	Pro	Ala	His
	Gly 545	Val	Thr	Ser	Ala	Pro 550	Asp	Asn	Arg	Pro	Ala 555	Leu	Gly	Ser	Thr	Ala 560
	Pro	Pro	Val	His	Asn 565	Val	Thr	Ser	Ala	Ser 570	Gly	Ser	Ala	Ser	Gly 575	Ser
	Ala	Ser	Thr	Leu 580	Val	His	Asn	GJA	Thr 585	Ser	Ala	Arg	Ala	Thr 590	Thr	Thr
1	Pro	Ala	Ser 595	Lys	Ser	Thr	Pro	Phe 600	Ser	Ile	Pro	Ser	His 605	His	Ser	Asp
	Thr	Pro 610	Thr	Thr	Leu	Ala	Ser 615	His	Ser	Thr	Lys	Thr 620	Asp	Ala	Ser	Ser
	Thr 625	His	His	Ser	Ser	Val 630	Pro	Pro	Leu	Thr	Ser 635	Ser	Asn	His	Ser	Thr 640
	Ser	Pro	Gln	Leu	Ser 645	Thr	Gly	Val	Ser	Phe 650	Phe	Phe	Leu	Ser	Phe 655	His
	Ile	Ser	Asn	Leu 660	Gln	Phe	Asn	Ser	Ser 665	Leu	Glu	Asp	Pro	Ser 670	Thr	Asp
	Tyr	Tyr	Gln 675	Glu	Leu	Gln	Arg	Asp 680	Ile	Ser	Glu	Met	Phe 685	Leu	Gln	Ile
	Tyr	Lys 690	Gln	Gly	Gly	Phe	Leu 695		Leu	Ser	Asn	Ile 700	_	Phe	Arg	Pro
	Gly 705	Ser	Val	Val	Val	Gln 710		Thr	Leu	Ala	Phe 715	Arg	Glu	Gly	Thr	Ile 720
	Asn	Val	His	Asp	Val 725	Glu	Thr	Gln	Phe	Asn 730		Туг	Lys	Thr	Glu 735	Ala

<210> 42 <211> 751 <212> DNA

<213> Homo sapiens

<400> 42

acgggcacgg ccggtaccat caatgtccac gacgtggaga cacagttcaa tcagtataaa 60 acggaagcag cctctcgata taacctgacg atctcagacg tcagcgtgag tgatgtgcca 120 ttteetttet etgeeeagte tggggetggg gtgceagget ggggeatege getgetggtg 180 ctggtctgtg ttctggttgc gctggccatt gtctatctca ttgccttggc tgtctgtcag 240 300 tgccgccgaa agaactacgg gcagctggac atctttccag cccgggatac ctaccatcct atgagogagt accocaccta ccacacccat gggogotatg tgccccctag cagtaccgat 360 cgtagcccct atgagaaggt ttctgcaggt aacggtggca gcagcctctc ttacacaaac 420 480 ccagcagtgg cagccgcttc tqccaacttg taggqcacgt cqccgctgag ctgagtqgcc agccagtgcc attccactcc actcaggttc ttcaggccag agcccctgca ccctgtttgg 540 600 gctggtgagc tgggagttca ggtgggctgc tcacagcctc cttcagaggc cccaccaatt 660 totoggacac ttotcagtgt gtggaagetc atgtgggccc ctgaggctca tgcctgggaa gtgttgtggg ggctcccagg aggactggcc cagagagccc tgagatagcg gggatcctga 720 751 actggactga ataaaacgtg gtctcccact g

<210> 43 <211> 820 <212> DNA <213> Homo sapiens

<400> 43

acggccggtt ttctgggcct ctccaatatt aagttcaggc caggatctgt ggtggtacaa 60 ttgactctgg ccttccgaga aggtaccatc aatgtccacq acgtggagac acagttcaat 120 cagtataaaa cggaagcagc ctctcgatat aacctgacga tctcagacgt cagcgtgagt 180 gatgtgccat ttcctttctc tgcccagtct ggggctgggg tgccaggctg gggcatcgcg 240 300 ctgctggtgc tggtctgtgt tctggttgcg ctggccattg tctatctcat tgccttggct gtctgtcagt gccgccgaaa gaactacggg cagctggaca tctttccagc ccgggatacc 360 taccatecta tgagegagta ecceaectae cacaeccatg ggegetatgt geceectage 420 agtaccgatc gtagccccta tgagaaggtt tctgcaggta acggtggcag cagcctctct 480 tacacaaacc cagcagtggc agccgcttct gccaacttgt agggcacgtc gccgctgagc 540 tgagtggcca gccagtgcca ttccactcca ctcaggttct tcaggccaga gcccctgcac 600 cctgtttggg ctggtgagct gggagttcag gtgggctgct cacagcctcc ttcagaggcc 660 ccaccaattt ctcggacact tctcagtgtg tggaagctca tgtgggcccc tgaggctcat 720 780 gcctgggaag tgttgtgggg gctcccagga ggactggccc agagagccct gagatagcgg ggatcctgaa ctggactgaa taaaacgtgg tctcccactg 820

<210> 44 <211> 1132 <212> DNA <213> Homo sapiens

<400> 44 acggccgcta ccacaaccc agccagcaag agcactccat totcaattcc cagccaccac 60 totgatactc ctaccaccct tgccagccat agcaccaaga ctgatgccag tagcactcac 120 catagetegg tacetectet cacetectee aateacagea ettetececa gttgtetaet 180 240 ggggtetett tettttteet gtetttteae attteaaace teeagtttaa tteetetetg gaagateeca geacegaeta etaceaagag etgeagagag acatttetga aatgtttttg 300 360 cagatttata aacaaggggg ttttctgggc ctctccaata ttaagttcag gccaggatct gtggtggtac aattgactct ggccttccga gaaggtacca tcaatgtcca cgacgtggag 420 acacagttca atcagtataa aacggaagca gcctctcgat ataacctgac gatctcagac 480 540 gtcagcgtga gtgatgtgcc atttcctttc tctgcccagt ctggggctgg ggtgccaggc 600 tggggcatcg cgctgctggt gctggtctgt gttctggttg cgctggccat tgtctatctc attgccttgg ctgtctgtca gtgccgccga aagaactacg ggcagctgga catctttcca 660

23/32	
gcccgggata cctaccatcc tatgagcgag taccccacct accacacca tgggcgctat	720
gtgcccccta gcagtaccga tcgtagcccc tatgagaagg tttctgcagg taacggtggc	780
agcagcctct cttacacaaa cccagcagtg gcagccgctt ctgccaactt gtagggcacg	840
tegeegetga getgagtgge eageeagtge catteeaete cacteaggtt etteaggeea	900
gagcccctgc accctgtttg ggctggtgag ctgggagttc aggtgggctg ctcacagcct	960
cetteagagg ceecaccaat tteteggaca etteteagtg tgtggaaget catgtgggee	1020
cctgaggctc atgcctggga agtgttgtgg gggctcccag gaggactggc ccagagagcc	1080
ctgagatagc ggggatcctg aactggactg aataaaacgt ggtctcccac tg	1132
<210> 45 <211> 717 <212> DNA <213> Homo sapiens <400> 45	
acaggttctg gtcatgcaag ctctacccca ggtggagaaa aggagacttc ggctacccag	60
agaagttcag tgcccagctc tactgagaag aatgctttta attcctctct ggaagatccc	120
agcaccgact actaccaaga gctgcagaga gacatttctg aaatgttttt gcagatttat	180
aaacaagggg gttttctggg cctctccaat attaagttca ggccaggatc tgtggtggta	240
caattgactc tggccttccg agaaggtacc atcaatgtcc acgacgtgga gacacagttc	300
aatcagtata aaacggaagc agcctctcga tataacctga cgatctcaga cgtcagcgtg	360
agtgatgtgc catttecttt etetgeceag tetggggetg gggtgceagg etggggeate	420
gegetgetgg tgetggtetg tgttetggtt gegetggeea ttgtetatet eattgeettg	480
gctgtctgtc agtgccgccg aaagaactac gggcagctgg acatctttcc agcccgggat	540
acctaccatc ctatgagega gtaccecace taccacacec atgggegeta tgtgeecect	600
agcagtaccg atcgtagccc ctatgagaag gtttctgcag gtaatggtgg cagcagcctc	660
tettacacaa acceageagt ggeageeact tetgeeaact tgtaggggea egtegee	717
<210> 46 <211> 2487 <212> DNA <213> Homo sapiens	
<400> 46 ctcgacccac gcgtccgctc gacccacgcg tccgcacctc ggccccggac accaggccgg	60
coccyggete caccyccec ccageccacy gtgtcacete ggccccggae accaggecgg	120
ccccgggctc caccgccccc ccagcccacg gtgtcacctc ggccccggac accaggccgg	180
ccccgggctc caccgccccc ccagcccacg gtgtcacctc ggccccggac accaggccgg	240

WO 2005/019269 PCT/US2004/027954 24/32

ccccgggctc	caccgccccc	ccagcccacg	gtgtcacctc	ggccccggac	accaggccgg	300
ccccgggctc	caccgccccc	ccagcccacg	gtgtcacctc	ggccccggac	accaggccgg	360
ccccgggctc	caccgccccc	ccagcccacg	gtgtcacctc	ggccccggac	accaggccgg	420
ccccgggctc	caecgeceee	ccagcccacg	gtgtcacctc	ggccccggac	accaggccgg	480
ccccgggctc	caccgccccc	ccagcccacg	gtgtcacctc	ggccccggac	accaggccgg	540
ccccgggctc	caccgccccc	ccagcccacg	gtgtcacctc	ggccccggac	accaggccgg	600
ccccgggctc	caccgccccc	ccagcccacg	gtgtcacctc	ggccccggac	accaggccgg	660
ccccgggctc	caccgccccc	ccagcccacg	gtgtcacctc	ggccccggac	accaggccgg	720
ccccgggctc	caccgccccc	ccagcccacg	gtgtcacctc	ggccccggac	accaggccgg	780
ccccgggctc	caccgccccc	ccagcccacg	gtgtcacctc	ggccccggac	accaggccgg	840
ccccgggctc	caccgccccc	ccagcccacg	gtgtcacctc	ggccccggac	accaggccgg	900
ccccgggctc	caccgccccc	ccagcccacg	gtgtcacctc	ggccccggac	accaggccgg	960
ccccgggctc	caccgccccc	ccagcccacg	gtgtcacctc	ggccccggac	accaggccgg	1020
ccccgggctc	caccgccccc	ccagcccacg	gtgtcacctc	ggccccggac	accaggccgg	1080
ccccgggctc	caccgccccc	ccagcccacg	gtgtcacctc	ggccccggac	accaggccgg	1140
ccccgggctc	caccgccccc	ccagcccacg	gtgtcacctc	ggccccggac	accaggccgg	1200
ccccgggctc	caccgccccc	ccagcccatg	gtgtcacctc	ggccccggac	aacaggcccg	1260
cettgggete	caccgcccct	ccagtccaca	atgtcacctc	ggcctcaggc	tctgcatcag	1320
geteagette	tactctggtg	cacaacggca	cctctgccag	ggctaccaca	accccagcca	1380
gcaagagcac	tccattctca	attcccagcc	accactctga	tactcctacc	acccttgcca	1440
gccatagcac	caagactgat	gccagtagca	ctcaccatag	ctcggtacct	cctctcacct	1500
cctccaatca	cagcacttct	ccccagttgt	ctactggggt	ctctttcttt	ttcctgtctt	1560
ttcacatttc	aaacctccag	tttaattcct	ctctggaaga	tcccagcacc	gactactacc	1620
aagagctgca	gagagacatt	tctgaaatgt	ttttgcagat	ttataaacaa	gggggttttc	1680
tgggcctctc	caatattaag	ttcaggccag	gatctgtggt	ggtacaattg	actctggcct	1740
tccgagaagg	taccatcaat	gtccacgacg	tggagacaca	gttcaatcag	tataaaacgg	1800
aagcagcctc	: tcgatataac	ctgacgatct	cagacgtcag	cgtgagtgat	gtgccatttc	1860
ctttctctgc	ccagtctggg	gctggggtgc	: caggctgggg	categegetg	ctggtgctgg	1920
tctgtgttct	ggttgegetg	gccattgtct	atctcattgo	cttggctgtc	tgtcagtgcc	1980
gccgaaagaa	ctacgggcag	ctggacatct	ttccagcccg	ggatacctac	catcctatga	2040

gcgagtaccc	cacctaccac	acccatgggc	gctatgtgcc	ccctagcagt	accgatcgta	2100
gcccctatga	gaaggtttct	gcaggtaacg	gtggcagcag	cctctcttac	acaaacccag	2160
cagtggcagc	cgcttctgcc	aacttgtagg	gcacgtcgcc	gctgagctga	gtggccagcc	2220
agtgccattc	cactccactc	aggttcttca	ggccagagcc	cctgcaccct	gtttgggctg	2280
gtgagctggg	agttcaggtg	ggctgctcac	agcctccttc	agaggcccca	ccaatttctc	2340
ggacacttct	cagtgtgtgg	aagctcatgt	gggcccctga	ggctcatgcc	tgggaagtgt	2400
tgtgggggct	cccaggagga	ctggcccaga	gagccctgag	atagcgggga	tcctgaactg	2460
gactgaataa	aacgtggtct	cccactg				2487

<210> 47

<211> 19

<212> PRT <213> Homo sapiens

. <400> 47

Met Thr Pro Gly Thr Gln Ser Pro Phe Phe Leu Leu Leu Leu Thr 10

Val Leu Thr

<210> 48

<211> 4003

<212> DNA

<213> Homo sapiens

<400> 48

acaggttctg gtcatgcaag ctctacccca ggtggagaaa aggagacttc ggctacccag 60 agaagttcag tgcccagctc tactgagaag aatgctgtga gtatgaccag cagcgtactc 120 tecagecaca geologistic aggetected accaeteagg gacaggatgt caetetggee 180 ceggecaegg aaccagette aggtteaget gecaeetggg gacaggatgt caeeteggte 240 ccagtcacca ggccagccct gggctccacc accccgccag cccacgatgt cacctcagcc 300 ccggacaaca agccagcccc gggctccacc gccccccag cccacggtgt cacctcggcc 360 420 · ccggacacca ggccggcccc gggctccacc gccccccag cccacggtgt cacctcggcc ccggacacca ggccggcccc gggctccacc gccccccag cccacggtgt cacctcggcc 480 coggacacca ggccggcccc gggctccacc gccccccag cccacggtgt cacctcggcc 540 600 coggacacca ggccggcccc gggctccacc gccccccag cccacggtgt cacctcggcc ceggacaeca ggceggeeee gggetecaee geceeecag eccaeggtgt caeeteggee 660 coggacacca ggccggcccc gggetccacc gccccccag cccacggtgt cacctcggcc 720

WO 2005/019269 PCT/US2004/027954 26/32

ccggacacca	ggccggcccc	gggctccacc	gccccccag	cccacggtgt	cacctcggcc	780
ccggacacca	ggccggcccc	gggctccacc	gccccccag	cccacggtgt	cacctcggcc	840
ccggacacca	ggccggcccc	gggctccacc	gccccccag	cccacggtgt	cacctcggcc	900
ccggacacca	ggccggcccc	gggctccacc	gccccccag	cccacggtgt	caccteggee	960
ccggacacca	ggccggcccc	gggctccacc	gccccccag	cccacggtgt	cacctcggcc	1020
ccggacacca	ggccggcccc	gggctccacc	gccccccag	cccacggtgt	caccteggee	1080
ccggacacca	ggccggcccc	gggctccacc	gccccccag	cccacggtgt	cacctcggcc	1140
ccggacacca	ggccggcccc	gggctccacc	gccccccag	cccacggtgt	cacctcggcc	1200
ccggacacca	ggccggcccc	gggctccacc	gcccccccag	cccacggtgt	caccteggee	1260
ccggacacca	gaccaacccc	gggctccacc	gcccccceag	cccacggtgt	caccteggee	1320
ccggacacca	ggeeggeeee	gggctccacc	gcccccccag	cccacggtgt	caccteggee	1380
ccggacacca	ggccggcccc	gggctccacc	gccccccag	cccacggtgt	cacctcggcc	1440
ccggacacca	ggccggcccc	gggctccace	gcccccccag	cccacggtgt	cacctcggcc	1500
ccggacacca	ggccggcccc	gggctccacc	gccccccag	cccacggtgt	caccteggee	1560
ccggacacca	ggccggcccc	gggctccacc	gcccccccag	cccacggtgt	cacctcggcc	1620
ccggacacca	ggccggcccc	gggctccacc	gccccccag	cccacggtgt	caccteggee	1680
ccggacacca	ggccggcccc	gggctccacc	gccccccag	cccacggtgt	cacctcggcc	1740
ccggacacca	ggccggcccc	gggctccacc	gccccccag	cccacggtgt	cacctcggcc	1800
ccggacacca	ggccggcccc	gggctccacc	gcccccccag	cccacggtgt	cacctcggcc	1860
ccggacacca	ggccggcccc	gggctccacc	gcccccccag	cccacggtgt	cacctcggcc	1920
ccggacacca	ggccggcccc	gggctccacc	gcccccccag	cccacggtgt	cacctcggcc	1980
ccggacacca	ggccggcccc	gggctccacc	gcccccccag	cccacggtgt	cacctcggcc	2040
ccggacacca	ggccggcccc	gggctccacc	gcccccccag	cccacggtgt	cacctcggcc	2100
ccggacacca	ggccggcccc	gggctccacc	gccccccag	cccacggtgt	cacctcggcc	2160
ccggacacca	ggccggcccc	gggctccacc	gcccccccag	cccacggtgt	cacctcggcc	2220
ccggacacca	ggccggcccc	gggctccacc	gcccccccag	cccacggtgt	cacctcggcc	2280
ccggacacca	ggccggcccc	gggctccacc	gcccccccag	cccacggtgt	cacctcggcc	2340
ccggacacca	ggccggcccc	gggctccacc	gcccccccag	cccacggtgt	caccteggee	2400
ccggacacca	ggccggcccc	gggctccacc	gcccccccag	cccacggtgt	cacctcggcc	2460
ccggacacca	ggccggcccc	gggctccacc	gcccccccag	cccacggtgt	cacctcggcc	2520
ccggacacca	ggccggccc	gggctccacc	gccccccag	cccacggtgt	cacctcggcc	2580

ccggacacca	ggccggcccc	gggctccacc	gccccccag	cccacggtgt	cacctcggcc	2640
ccggacacca	ggccggcccc	gggctccacc	gccccccag	cccacggtgt	cacctcggcc	2700
ccggacacca	ggccggcccc	gggctccacc	gccccccag	cccatggtgt	cacctcggcc	2760
ccggacaaca	ggcccgcctt	gggctccacc	gcccctccag	tccacaatgt	cacctcggcc	2820
tcaggctctg	catcaggete	agcttctact	ctggtgcaca	acggcacctc	tgccagggct	2880
accacaaccc	cagccagcaa	gagcactcca	ttctcaattc	ccagccacca	ctctgatact	2940
cctaccaccc	ttgccagcca	tagcaccaag	actgatgcca	gtagcactca	ccatagctcg	3000
gtacctcctc	tcacctcctc	caatcacage	acttctcccc	agttgtctac	tggggtctct	3060
ttetttttce	tgtcttttca	catttcaaac	ctccagttta	attcctctct	ggaagatccc	3120
agcaccgact	actaccaaga	gctgcagaga	gacatttctg	aaatgttttt	gcagatttat	3180
aaacaagggg	gttttctggg	cctctccaat	attaagttca	ggccaggatc	tgtggtggta	3240
caattgactc	tggccttccg	agaaggtacc	atcaatgtcc	acgacgtgga	gacacagttc	3300
aatcagtata	aaacggaagc	agcctctcga	tataacctga	cgatctcaga	cgtcagcgtg	3360
agtgatgtgc	catttccttt	ctctgcccag	tctggggctg	gggtgccagg	ctggggcatc	3420
gcgctgctgg	tgctggtctg	tgttctggtt	gcgctggcca	ttgtctatct	cattgccttg	3480
gctgtctgtc	agtgccgccg	aaagaactac	gggcagctgg	acatctttcc	agcccgggat	3540
acctaccatc	ctatgagcga	gtaccccacc	taccacaccc	atgggcgcta	tgtgccccct	3600
agcagtaccg	ategtagece	ctatgagaag	gtttctgcag	gtaacggtgg	cagcagcctc	3660
tcttacacaa	acccagcagt	ggcagccgct	tctgccaact	tgtagggcac	gtcgccgctg	3720
agctgagtgg	, ccagccagtg	ccattccact	. ccactcaggt	tcttcaggcc	agagcccctg	3780
caccctgttt	: gggctggtga	gctgggagtt	: caggtgggct	gctcacaged	tccttcagag	3840
gccccaccaa	tttctcggac	acttctcagt	gtgtggaago	: tcatgtgggc	ccctgaggct	3900
catgcctggg	g aagtgttgtg	ggggctccca	a ggaggactgg	cccagagagc	cctgagatag	3960
cggggatcct	gaactggact	gaataaaac	g tggtctccca	ctg		4003

<220>

<223> PCR Primer

<400> 49

gggaattcat gacaccgggc acccagtc

<210> 49 <211> 28 <212> DNA <213> Artificial Sequence

<210> <211> <212> <213>	50 27 DNA Artificial Sequence	
<220>		
<223>	PCR Primer	
<400> ggtctc	50 gaga acaactgtaa gcactgt	27
<210> <211>		
<212>		
<220>	•	
<223>	PCR Primer	
<400> ggtcgg	51 ccgt aacaactgta agcactgt	28
<210> <211> <212> <213>	28 .	
<220>		
<223>	PCR Primer	
<400> gcacgo	52 geoge taccacaacc ccagccag	28
<210> <211> <212> <213>	28 DNA	
<220>		
<223>	PCR Primer	
<400> gcacgo	53 geogg ttttetggge etetecaa	28
<210> <211> <212> <213>	29	
<220>		
<223>	PCR Primer	

<400> 54 gcacggccgg taccatcaat gtccacgac	29 .
<210> 55 <211> 28 <212> DNA <213> Artificial Sequence	
<220>	
<223> PCR Primer	
<400> 55 gggggatcct acaagttggc agaagcgg	28
<210> 56 <211> 39 <212> DNA <213> Artificial Sequence	
<220>	
<223> PCR Primer	
<400> 56 tgctccac agtgcttaca ggttctggtc atgcaagct	39
<210> 57 <211> 32 <212> DNA <213> Artificial Sequence	
<223> PCR Primer	
<400> 57 gagcttgcat gaccagaacc tgtaacaact gt	32 .
<210> 58 <211> 23 <212> PRT <213> Homo sapiens	
<400> 58	
Met Thr Pro Gly Thr Gln Ser Pro Phe Phe Leu Leu Leu Leu Thr 1 5 10 15	
Val Leu Thr Val Val Thr Ala 20	
<210> 59 <211> 24 <212> PRT	

<213> Homo sapiens

<400> 59

Met Thr Pro Gly Thr Gln Ser Pro Phe Phe Leu Leu Leu Leu Thr

Val Leu Thr Val Val Thr Ala Gly

<210> 60

<211> 50 <212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Peptide

<400> 60

Thr Ile Asn Val His Asp Val Glu Thr Gln Phe Asn Gln Tyr Lys Thr

Glu Ala Ala Ser Pro Tyr Asn Leu Thr Ile Ser Asp Val Ser Val Ser

Asp Val Pro Phe Pro Phe Ser Ala Gln Ser Gly Ala His His His His 40

His His 50

<210> 61 <211> 63 <212> PRT <213> Artificial Sequence

<220>

<223> Synthetic Peptide

<400> 61

Ser Val Val Gln Leu Thr Leu Ala Phe Arg Glu Gly Thr Ile Asn

Val His Asp Val Glu Thr Gln Phe Asn Gln Tyr Lys Thr Glu Ala Ala

Ser Pro Tyr Asn Leu Thr Ile Ser Asp Val Ser Val Ser Asp Val Pro

Phe Pro Phe Ser Ala Gln Ser Gly Ala His His His His His

<210> 62

<211> 19

<212> PRT <213> Artificial Sequence

<220>

<223> Synthetic Peptide

<400> 62

His His His His His Ser Val Val Gln Leu Thr Leu Ala Phe

Arg Glu Gly

<210> 63 <211> 44

<212> PRT

<213> Homo sapiens

<400> 63

Thr Ile Asn Val His Asp Val Glu Thr Gln Phe Asn Gln Tyr Lys Thr

Glu Ala Ala Ser Arg Tyr Asn Leu Thr Ile Ser Asp Val Ser Val Ser

Asp Val Pro Phe Pro Phe Ser Ala Gln Ser Gly Ala

<210> 64

<211> 44

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic Peptide

<400> 64

Thr Ile Asn Val His Asp Val Glu Thr Gln Phe Asn Gln Tyr Lys Thr

Glu Ala Ala Ser Pro Tyr Asn Leu Thr Ile Ser Asp Val Ser Val Ser

Asp Val Pro Phe Pro Phe Ser Ala Gln Ser Gly Ala

<210> 65

<211> 13

2

<212> PRT

<213> Homo sapiens

<400> 65

Ser Val Val Gln Leu Thr Leu Ala Phe Arg Glu Gly

<210> 66

<211> 57 <212> PRT <213> Homo sapiens

<400> 66

Ser Val Val Val Gln Leu Thr Leu Ala Phe Arg Glu Gly Thr Ile Asn

Val His Asp Val Glu Thr Gln Phe Asn Gln Tyr Lys Thr Glu Ala Ala

Ser Pro Tyr Asn Leu Thr Ile Ser Asp Val Ser Val Ser Asp Val Pro 40

Phe Pro Phe Ser Ala Gln Ser Gly Ala